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Metabotyping for the development of tailored dietary advice solutions in a European population: the Food4Me study

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Shortened title

Tailored advice for a European population

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Abstract

Traditionally, personalised nutrition was delivered at an individual level. However, the concept of delivering tailored dietary advice at a group level through the identification of metabotypes or groups of metabolically similar individuals has emerged. Whilst this approach to personalised nutrition looks promising, further work is needed to examine this concept across a wider population group. Therefore, the objectives of this study are to 1) identify metabotypes in a European population and 2) develop targeted dietary advice solutions for these metabotypes. Using data from the Food4Me study (n = 1,607), k-means cluster analysis revealed the presence of three metabolically distinct clusters based on twenty-seven metabolic markers including cholesterol, individual fatty acids and carotenoids. Cluster 2 was identified as a metabolically healthy metabotype as these individuals had the highest omega 3 index ($6.56 \pm 1.29 \%$), carotenoids ($2.15 \pm 0.71 \mu\text{M}$) and lowest total saturated fat levels. Based on its fatty acid profile, cluster 1 was characterised as a metabolically unhealthy cluster. Targeted dietary advice solutions were developed per cluster using a decision tree approach. Testing of the approach was performed by comparison with the personalised dietary advice, delivered by nutritionists, to Food4Me study participants (n = 180). Excellent agreement was observed between the targeted and individualised approaches with an average match of 82 % at the level of delivery of the same dietary message. Future work should ascertain whether this proposed method could be utilised in a healthcare setting, for the rapid and efficient delivery of tailored dietary advice solutions.

Introduction

Early definitions of personalised nutrition were gene focused, however, in recent times, the definition has been extended and now incorporates the concept of levels⁽¹⁾. This reworked definition of personalised nutrition now includes Level 1 personalised advice which involves delivering personalised advice based on dietary intake, Level 2 personalised advice which involves personalised advice based on diet and phenotypic markers such as blood markers and BMI, and Level 3 personalised advice which builds on the previous levels and includes diet, phenotype and genotype information⁽²⁾. Whilst such definitions focus on personalised advice delivered at an individual level, there is an emerging concept that has gained momentum in recent years, where dietary advice can be tailored to specific groups of individuals and is referred to as targeted nutrition^(3; 4; 5).

These groups of individuals have similar characteristics and are referred to as metabotypes⁽⁶⁾. There are numerous examples of metabotyping in the medical literature where it has been utilised to sub-group patients with diseases with differential symptomology^(7; 8; 9; 10). For example, several studies have used cluster analysis to identify sub-groups of patients with characteristic phenotypes of asthma, a disease which is very heterogeneous in nature^(11; 12; 13; 14). Metabotyping has also been used to identify groups of individuals with differing responses to drug treatments^(15; 16; 17) and dietary interventions^(18; 19).

However, while there are many examples of identifying groups of similar individuals in the population^(7; 8; 9; 20), the evidence base for developing tailored health solutions for these groups is weak. Previous work from our group demonstrated a framework for the delivery of targeted nutrition advice to metabolically similar groups or metabotypes in the population⁽²¹⁾. In this study, three distinctly different metabotypes were identified on the basis of four routinely measured markers of metabolic health including blood triacylglycerols, total cholesterol, HDL cholesterol and glucose (n=896). Using a decision tree approach, targeted dietary advice messages were developed based on the characteristics of each cluster. Good agreement was observed between the targeted dietary advice method and an individualised method without the need for collection of detailed dietary data⁽²¹⁾. Overall, this previous work demonstrated the potential of the metabotyping approach to deliver appropriate tailored dietary advice at a group level with minimal data collection required.

In the current study, this concept is further advanced using data from the Food4Me study, a personalised nutrition intervention study⁽²²⁾. In Food4Me, participants received personalised advice based on the three levels of personalisation, delivered by trained nutritionists and thus provides a valuable resource for testing the targeted nutrition approach⁽²²⁾. Therefore, the objectives of this study were to 1) identify metabotypes in a European population group and 2) develop and test targeted dietary advice solutions for these metabotypes by comparison with the personalised dietary advice given within the Food4Me study.

Materials and Methods

Study design and ethical approval

As part of the Food4Me project (ClinicalTrials.gov number: NCT01530139, <https://clinicaltrials.gov/ct2/show/NCT01530139>), a proof-of-principle (PoP) study was conducted, which compared the effectiveness of personalised nutrition advice, based on the three levels of personalisation, on health related outcomes, compared with generic healthy eating advice. This was an internet-based study, designed to emulate a personalised nutrition service, and was conducted in seven research centres across Europe. Ethical approval was obtained from the Research Ethics Committees at each university or research centre. This study was conducted according to the guidelines laid down in the Declaration of Helsinki and all procedures involving human subjects were approved by the Research Ethics Committees at each university or research centre. Participants (n = 1,607) were randomised into one of four groups; Control group which received general European based healthy eating guidelines, Level 1 participants received personalised advice based on their dietary intake, Level 2 received personalised advice based on their diet and phenotype and Level 3 received advice on their diet, phenotype and genotype. More details on the overall study design can be found elsewhere^(22; 23). Written informed consent was obtained from all subjects.

Data collection and personalised feedback

All data were self-collected by participants using detailed instructions provided by researchers and online video demonstrations. A more detailed description of the data collection methods is reported elsewhere⁽²²⁾. In brief, habitual dietary intake was assessed using the online Food4Me food frequency questionnaire (FFQ), which was previously developed and validated for the purposes of the study^(24; 25). The foods included in the FFQ were aggregated to form thirty-two food groups. The list of the foods contributing to each of the food groups is found in

Supplementary Table 1. Participants were provided with a measuring tape to perform anthropometric measures including weight (kg), height (m) and circumferences including waist (cm), hip (cm) and thigh (cm); all were collected according to standard previously published protocols⁽²²⁾. A validation study was conducted to assess the accuracy of these measurements and strong correlation coefficients were observed between the self-reported and measurements performed face-to-face by researchers⁽²⁶⁾.

Metabolic markers were measured by finger-prick blood samples collected by participants using a collection pack provided by Vitas Ltd (Oslo, Norway). Participants were asked to fast 8 hours prior to collection in the morning and filled two dry blood spot (DBS) cards (five drops of blood or 150µl of blood per card). Once filled, cards were left to dry for 2-4 hours at room temperature and placed in an airtight aluminium bag with a drying sachet and returned by post to their corresponding recruiting centre. The samples were then sent via courier service to Vitas, where the following metabolic markers were measured: total cholesterol, carotenoids (lutein, zeaxanthin, beta-cryptoxanthin, alpha-caroten, beta-carotene, lycopene) and twenty fatty acids as shown in **Table 1**. The metabolic markers were measured using the following methods: cholesterol (LC-UV), carotenoids (HPLC-DAD-MS/MS) and fatty acids (GC-FID).

Participants randomised to levels 1, 2 and 3 received personalised reports based on decision trees to allow for the delivery of systematic tailored advice. The personalised reports were sent via email at months 0, 3 and 6. Standard operating procedures were developed for use of the decision trees and these were standardised across the seven recruitment centres to ensure consistency in the personalised advice given across all centres. Those individuals in Level 1 received feedback based on their current dietary intake and physical activity levels. Level 2 participants received feedback on based on their current diet, physical activity levels and phenotypic measures such as anthropometry and metabolic markers. Level 3 participants received the same feedback as Level 2 with the addition of genotypic information. The final section of the report contained a personalised goals section where participants were given three nutrient-related goals. The personalised goals were selected by a pre-defined ranking system, where those nutrients and metabolic markers that most warranted change, were prioritised. Participants were asked to focus on making changes to these three nutrients in the personalised reports in line with the patient-centred counselling models for facilitating behaviour change⁽²⁷⁾.

Statistics

Baseline data were analysed using SPSS software package version 20.0 for Windows (SPSS, Inc. Chicago, IL, USA). Twenty-seven metabolic markers including cholesterol and individual carotenoids and fatty acids from the DBS analysis were chosen for clustering as presented in **Table 1**. Following standardisation using z-scores, two-step cluster analysis revealed the presence of three clusters and k-means cluster analysis was then used to characterise the clusters. The differences between the clusters were assessed using one-way ANOVA with Bonferroni post-hoc tests. Chi-square distributions were used to assess categorical variables across the clusters including gender and country. As age, gender, BMI and country were significantly different across the clusters, these variables were controlled for in the general linear models with Bonferroni post hoc tests. P values were also adjusted for multiple comparisons using the Bonferroni approach.

Development and testing of targeted dietary advice

Targeted dietary advice was developed for each cluster based on the characteristics of the cluster and using a decision tree process. Two decision trees were developed per cluster based on: 1) metabolic markers and anthropometric information and 2) dietary information. This resulted in forty-nine messages for cluster 1, twenty messages for cluster 2 and twenty-four messages for cluster 3. The cut-offs used for the metabolic markers, anthropometric and dietary data within the decision trees are presented in **Table 5**. Since there are no defined cut-offs for total saturated fat (%) from DBS data, cluster 1 was described as high saturated fat, cluster 2 low saturated fat and cluster 3 medium saturated fat based on the mean values across the clusters as shown in **Table 1**.

The appropriateness of the targeted dietary advice developed per cluster was then tested by comparison with the three nutrient-related goals, that were delivered to all of Level 2 participants (n = 180) by trained nutritionists, as part of their personalised feedback reports. The agreement between the two methods was assessed based on the following questions:

1. How many of the nutrient-related goals given as part of the personalised advice reports within the Food4Me study were given as part of the targeted dietary advice derived from this study?
2. How many dietary messages were given as part of the targeted dietary advice in comparison with the personalised advice within Food4Me? i.e. number of messages given as per the targeted dietary advice.

Results

Characterisation of the clusters

Three clusters were identified in the Food4Me population (**Table 1**). Cluster 1 ($n = 326$) was the group with the highest cholesterol, highest circulating trans fatty acids ($0.85 \pm 0.25 \%$) and lowest omega-3 index ($5.16 \pm 0.93 \%$). Cluster 2 ($n = 433$) was the most metabolically healthy group as they had the highest average omega-3 index ($6.56 \pm 1.29 \%$), highest total carotenoid concentrations ($2.15 \pm 0.71 \mu\text{M}$) and lowest total saturated fat. Cluster 3 subjects ($n = 595$) had the lowest average cholesterol concentrations ($4.25 \pm 0.78 \text{ mM}$) and highest stearic acid (**Table 1**). Age was significantly different across the groups with cluster 1 and 2 being older on average (**Table 2**). BMI and waist circumference were also significantly different across the clusters. Cluster 1 had the highest BMI of $27.7 \pm 5.3 \text{ kg/m}^2$ and waist circumference (0.93 ± 0.14

m) while participants in cluster 2 had the lowest BMI and waist circumference (**Table 2**). With the exception of the Netherlands and United Kingdom, the distribution of nationality differed significantly across the clusters.

Reported dietary intakes across the clusters are presented in **Table 3**. There were no differences in total energy intake and macronutrients across the clusters. However, percentage energy contribution from alcohol and polyunsaturated fatty acids were found to be significantly different ($p = 0.048$). Furthermore, intakes of many micronutrients differed significantly across the clusters including fat soluble vitamins A, D and E, as well as some water soluble vitamins such as folate, vitamin B6 and vitamin C. Participants in cluster 1 had the higher percentage contribution of energy from alcohol ($4.2 \pm 4.5 \%$) compared with individuals in cluster 2 and cluster 3. The diets of cluster 2 participants were considered to be healthier as these individuals had the highest intakes of dietary fibre ($32 \pm 15 \text{ g}$), fat soluble vitamins D and E, folate and vitamin C.

Intakes of the food groups savouries ($p = 1.27 \times 10^{-4}$), fruit ($p = 1.39 \times 10^{-8}$), fish, fish dishes and products ($p = 8.16 \times 10^{-4}$) differed significantly between the clusters as illustrated in **Table 4**. Similar to their nutrient intakes, participants in cluster 2 had the healthiest food intakes with the lowest intakes of savouries ($11 \pm 13 \text{ g}$) and white bread/rolls/scones/croissants ($34 \pm 73 \text{ g}$) and highest intakes of yoghurt ($91 \pm 107 \text{ g}$), fruit ($355 \pm 306 \text{ g}$), fish, fish dishes and products ($71 \pm 53 \text{ g}$). Clusters also differed in terms of supplement users ($p = 9.31 \times 10^{-8}$), with the highest percentage found in cluster 2 (54.3 %) who also had the highest omega-3 index.

Development of the targeted dietary advice

Targeted dietary advice was developed based on the characteristics (anthropometric, metabolic and nutrient intake data) of each cluster using a decision tree method. Two decision trees were constructed per cluster; a combined metabolic & anthropometric decision tree and a dietary decision tree. Ranges of the metabolic markers and nutrients were calculated for each of the clusters and these values were then used to determine whether individuals in each cluster were within the desirable or high/low range for that particular variable as shown in **Table 5**. The cut-offs used in the current study were based on those used within the Food4Me study (**Supplementary Table 2**), but were simplified for the purposes of the development of the targeted dietary advice. For the targeted dietary advice, the cut-offs were set as either 'desirable' or 'high/low' (**Table 5**), whereas in Food4Me the cut-offs were developed using a more complex gradation scale (**Supplementary Table 2**). This information was then used to construct the branches of each of the decision trees per cluster. Using this method, dietary advice was developed based on four metabolic markers (total cholesterol, total saturated fat, omega-3 index and carotenoids) and five key nutrients (salt, dietary fibre, iron, calcium and folate). Supplementary figures **1a**) and **1b**) demonstrate the metabolic and anthropometric decision tree and dietary decision trees for cluster 2 respectively and examples of a targeted message from each of the decision trees for cluster 2.

Comparison of the targeted dietary advice and personalised feedback reports

Level 2 participants from Food4Me (n = 180) were selected to test the appropriateness of the targeted dietary advice developed within this study. Excellent agreement was found between the personalised advice delivered by trained nutritionists in Food4Me and the targeted method developed in this study, with an average match of 82 % in relation to the dietary messages given (**Table 6**). Examining the clusters individually, good agreement was also found with an average match of 83 % for cluster 1, 74 % for cluster 2 and 88 % for cluster 3 for the dietary messages given. The number of messages given as part of the targeted dietary advice is depicted in **Table 7**. In general, more messages were given using the targeted approach compared with the individualised method used in Food4Me, where a restriction to three nutrient related goals was imposed.

Discussion

The present study demonstrates a successful method for the delivery of targeted nutrition advice using a combination of metabotyping and decision trees. Excellent agreement between

this method and that of a personalised method delivered by a team of trained nutritionists and dietitians in the Food4Me study was found, with an average match of 82 %, at the level of agreement of the same dietary message given. To the best of our knowledge, this is the first study to identify metabotypes in the European population and to develop tailored dietary solutions appropriate for participants from diverse cultures and dietary intakes. This work paves the way for further development of this approach and potential delivery of personalised nutrition advice to large population groups.

Using cluster analysis, three distinctly different metabotypes were identified based on a range of blood-based metabolic markers. Individuals in cluster 1 were found to have an unhealthy metabolic profile as these individuals had the highest cholesterol levels, highest saturated fat levels and lowest omega-3 index. On the other hand, individuals in cluster 2 was identified as the healthiest group and had the lowest saturated fat levels, highest carotenoid concentrations and highest omega-3 index. Subjects in cluster 3 were found to have the lowest cholesterol and carotenoid concentrations. These findings are similar to previously published studies on metabotypes^(6; 17). Morris and colleagues identified four metabotypes consisting of four different responses to an oral glucose tolerance test (OGTT)⁽⁶⁾. Classification of individuals based on their response curves to an OGTT revealed an ‘at-risk’ metabolic phenotype, which had the highest BMI, triacylglycerol levels, C-reactive protein, C-peptide, insulin and HOMA-IR score⁽⁶⁾. In a similar manner, van Bochove and colleagues identified three clusters based on their lipoprotein profiles and reported one cluster who did not respond favourably to fenofibrate treatment⁽¹⁷⁾. In our previous study, one cluster with a metabolically unfavourable profile and another cluster in which the individuals were relatively healthy with respect to a range of metabolic markers were also identified⁽²¹⁾. The consistency of identification of clusters across a range of studies adds validation to the approach and supports the clusters found in the present study.

An important finding from the current study is the evidence that there was a relationship between the metabolic profiles of each cluster and the corresponding nutrient and food group intakes of those clusters. For example, in line with their high carotenoid concentrations, participants in cluster 2 were also found to have the highest intakes of vitamin C, folate and dietary fibre. Similarly, individuals in cluster 2 had the highest intake of the fish, fish dishes and products which was also reflected in their metabolic profile as this group had the highest average omega-3 index. However, individuals in cluster 2 had the highest intakes of supplements which were likely to contribute to their high omega-3 levels. The agreement

between the metabolic profiles and dietary intake support the concept of using blood-based metabolotypes as a basis for targeted nutrition advice.

Good agreement between the proposed framework and the individualised advice delivered in the Food4Me study was observed. In Food4Me, personalised dietary advice was delivered by trained dietitians and nutritionists across seven research centres in Europe and was based on a decision tree method, which resulted in 295 possible dietary messages⁽²²⁾. In the final section of the personalised reports, participants were given three key pieces of dietary advice that they were encouraged to focus on, which were selected based a priority system, developed specially for the purposes of the Food4Me study⁽²²⁾. In contrast to this, a more simplified method is proposed here, in which blood-based metabolic data in conjunction with minimal dietary information could be used to deliver tailored dietary advice. This more simplified approach showed an average match of 82 % at the level of the dietary advice given, with the actual advice delivered within the Food4Me study. Based on this, it is proposed that tailored dietary advice could be given based primarily on the metabolic markers and information on the intakes of five key nutrients.

A framework for the delivery of targeted dietary advice in the Irish population, by the identification of three diverse metabolotypes, and development of tailored dietary advice based on decision trees was previously presented⁽²¹⁾. In the current paper, a similar method to identify metabolotypes was employed but we have advanced this concept by the inclusion of a broader range of metabolic data. Furthermore, the decision trees for the delivery of the advice were expanded to include specific key nutrients. This approach has potential to improve public health through the provision of tailored dietary advice to patients, in a quick and efficient manner, with minimal effort required by healthcare providers.

In this study, the metabolic markers were collected using DBS cards by the participants in their own homes. Collection of samples by DBS has a number of advantages including reduced costs, possibility of collection of large sample sizes, no blood processing and minimal storage facilities required^(28; 29). This presents another opportunity for the proposed framework to be adopted in the community setting where community health nurses could deliver the targeted dietary advice. Community nurses are suitable candidates to deliver tailored advice as they routinely see patients that may benefit from dietary/lifestyle change, have regular contact with patients over long periods, visit patients in their own homes and can involve their family in the intervention, and visit those who may not be physically capable of attending their doctor⁽³⁰⁾.

Chan and colleagues conducted a study to investigate the scope for risk management practices by nurses based in the community⁽³⁰⁾. They reported that levels of obesity and prevalence of risk factors including smoking status and low physical activity levels were higher in the individuals (n = 804) who took part in the study, compared with the general population, and that the majority of individuals with at least one risk factor had not received advice or been referred in the last three months⁽³⁰⁾. This suggests that there is considerable scope to deliver dietary and lifestyle interventions in the community. In addition, when provided with appropriate training, community nurses were shown to be confident in assessing lifestyle factors such as smoking, anthropometric measures and dietary intake⁽³¹⁾. It is envisaged that the proposed framework, in our study, could easily be adopted by nurses in the community setting, to deliver tailored dietary advice with minimal training required, and have the potential to reach many more individuals who could benefit from tailored dietary advice.

A major strength of this study is its applicability to the European population. Furthermore, good agreement was reported between the proposed targeted method and an individualised method delivered by a team of nutritionists across seven research centres in the Food4Me study. A limitation of this study is that the dietary intake data was collected using an online FFQ which assessed dietary intake of the previous month. Furthermore, the dietary advice developed did not take into account cooking abilities, likes/dislikes or cost of meals.

The present study developed a framework for the identification of metabotypes in the European population and the development of tailored dietary advice. Good agreement was found in comparison with an individualised personalised nutrition approach which has been used to deliver advice across seven countries. The demonstration of this approach in a pan European study offers significant credibility to the framework. In our previous study, we envisaged translation of this approach for use by healthcare professionals and the present study further supports such a concept. With this in mind, future work should test this framework in such a setting to ascertain whether the advice is effective in motivating changes in diet and lifestyle factors.

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Conflict of interest

None

Authorship

C.B.O.D., E.R.G. and L.B. developed and tested the targeted dietary approach, carried out the statistical analyses and drafted the manuscript. C.C.M., R.F., A.L.M., C.F.M.M., S.N.C., R.S.C., C.B.O.D, H.F., C.W., S.K., L.T., C.M., C.P.L., G.M., M.G., A.S., M.C.W. and J.C.M. conducted the intervention. I.T., C.A.D., H.D., Y.M., J.A.M., W.H.M.S., J.A.L., J.C.M. , M.J.G., E.R.G., and L.B. contributed to the research design of the Food4Me study. All authors contributed to a critical review of the manuscript during the writing process and approved the final version to be published.

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Table 1 Clustering variables and other metabolites

	Cluster 1 (N = 326)		Cluster 2 (N = 433)		Cluster 3 (N = 595)		
Clustering variables	Mean	SD	Mean	SD	Mean	SD	P value
Total Cholesterol (mM)	<u>5.01</u> ^{2,3}	1.02	4.79 ^{1,3}	0.91	4.25 ^{1,2}	0.78	1.30 x 10 ⁻³⁷
<i>Fatty acids (%)</i>							
Myristic acid C14:0	<u>1.07</u> ^{2,3}	0.52	0.69 ^{1,3}	0.24	0.59 ^{1,2}	0.20	5.06 x 10 ⁻⁹²
Pentadecyclic acid C15:0	0.22 ³	0.07	0.22 ³	0.06	0.18 ^{1,2}	0.04	1.03 x 10 ⁻³²
Palmitic acid C16:0	<u>24.77</u> ^{2,3}	1.87	22.62 ^{1,3}	1.48	22.94 ^{1,2}	1.45	1.39 x 10 ⁻⁷⁶
Palmitoleic acid C16:1	<u>1.82</u> ^{2,3}	0.56	1.16 ^{1,3}	0.37	1.02 ^{1,2}	0.34	2.22 x 10 ⁻¹³⁷
Margaric acid C17:0	0.30 ²	0.06	<u>0.34</u> ^{1,3}	0.06	0.31 ²	0.06	1.27 x 10 ⁻¹⁶
Stearic acid C18:0	12.07 ^{2,3}	1.12	12.79 ^{1,3}	1.00	<u>13.59</u> ^{1,2}	1.12	8.18 x 10 ⁻⁸²
cisVaccenic acid C18:1 cis	<u>1.52</u> ^{2,3}	0.32	1.42 ¹	0.25	1.43 ¹	0.23	1.47 x 10 ⁻⁷
Oleic acid C18:1	<u>20.72</u> ^{2,3}	2.4	18.06 ^{1,3}	1.65	18.80 ^{1,2}	1.86	1.02 x 10 ⁻⁷⁰
Arachidic acid C20:0	0.18 ^{2,3}	0.06	0.20 ^{1,3}	0.07	<u>0.23</u> ^{1,2}	0.09	1.02 x 10 ⁻¹⁹
Eicosenoic acid C20:1	0.25 ^{2,3}	0.06	0.27 ^{1,3}	0.06	<u>0.28</u> ^{1,2}	0.06	2.54 x 10 ⁻¹⁴
Total saturated fat*	<u>37.91</u> ^{2,3}	2.5	36.11 ^{1,3}	1.97	37.11 ^{1,2}	1.92	2.68 x 10 ⁻³⁰
Trans fatty acids	<u>0.85</u> ^{2,3}	0.25	0.79 ^{1,3}	0.24	0.75 ^{1,2}	0.21	1.07 x 10 ⁻¹⁰
Alphalinolenic acid C18:3 n3	<u>0.39</u> ³	0.19	0.37 ³	0.12	0.28 ^{1,2}	0.12	6.28 x 10 ⁻³⁹
Eicosapentaenoic acid C20:5 n3	0.66 ^{2,3}	0.35	<u>1.06</u> ^{1,3}	± 0.65	0.55 ^{1,2}	0.27	1.68 x 10 ⁻⁶⁷
Docosapentaenoic acid C22:5 n3	1.24 ^{2,3}	0.34	<u>1.56</u> ^{1,3}	± 0.33	1.35 ^{1,2}	0.37	2.81 x 10 ⁻³⁷
Docosahexaenoic acid C22:6 n3	2.57 ^{2,3}	0.72	<u>3.53</u> ^{1,3}	0.88	2.87 ^{1,2}	0.76	1.90 x 10 ⁻⁶⁰
Omega-3 index [†]	5.16 ^{2,3}	0.93	<u>6.56</u> ^{1,3}	1.29	5.41 ^{1,2}	0.92	5.14 x 10 ⁻⁸⁰

Linoleic acid C18:2 n6	17.55 ^{2,3}	2.09	<u>20.10</u> ^{1,3}	2.28	19.64 ^{1,2}	2.10	8.71 x 10 ⁻⁵⁸
Gamma-linolenic acid C18:3 n6	<u>0.24</u> ^{2,3}	0.10	0.16 ^{1,3}	0.07	0.18 ^{1,2}	0.07	1.10 x 10 ⁻³⁶
Eicosadienoic acid C20:2 n6	0.21 ^{2,3}	0.04	0.22 ^{1,3}	0.04	<u>0.24</u> ^{1,2}	0.04	8.04 x 10 ⁻³⁸
Dihomo-gamma-linolenic acid C20:3 n6	1.54 ^{2,3}	0.34	1.41 ^{1,3}	0.32	<u>1.60</u> ^{1,2}	0.33	6.00 x 10 ⁻¹⁹
Arachidonic acid C20:4 n6	7.93 ^{2,3}	1.46	8.64 ^{1,3}	1.23	<u>9.44</u> ^{1,2}	1.34	3.29 x 10 ⁻⁵⁶
<i>Carotenoids (μM)</i>							
aCaroten	0.08 ²	0.07	<u>0.21</u> ^{1,3}	0.17	0.08 ²	0.05	4.30 x 10 ⁻⁸⁴
bCaroten	0.28 ²	0.17	<u>0.66</u> ^{1,3}	0.36	0.27 ²	0.14	1.48 x 10 ⁻¹³²
bCryptoxanthin	0.14 ²	0.12	<u>0.29</u> ^{1,3}	0.22	0.16 ²	0.11	1.04 x 10 ⁻⁴⁸
Lutein	0.20 ^{2,3}	0.09	<u>0.29</u> ^{1,3}	0.15	0.18 ^{1,2}	0.08	4.56 x 10 ⁻⁵⁵
Lycopene	0.53 ²	0.24	<u>0.65</u> ^{1,3}	0.31	0.50 ²	0.23	7.68 x 10 ⁻¹⁹
Zeaxanthin	0.05 ²	0.03	<u>0.06</u> ^{1,3}	0.04	0.04 ²	0.03	1.12 x 10 ⁻¹⁸
Total carotenoids [‡]	1.28 ²	0.46	<u>2.15</u> ^{1,3}	0.71	1.21 ²	0.40	1.90 x 10 ⁻¹⁴⁵

N, number of participants. *Total saturated fat was calculated as the sum of myristic acid (C14:0), palmitic acid (C16:0) and stearic acid (C18:0). [†]Omega-3 index was calculated by the formula: omega-3 index = 1.4473 + 0.8303*(EPA+DPA+DHA). [‡]Total carotenoids was calculated by the following formula: Total carotenoids = alpha-carotene + beta-carotene + lutein + zeaxanthin + beta-cryptoxanthin + lycopene. Values are presented as means ± standard deviations. One-way ANOVA was used to examine the differences across the clusters. Underlined values indicate the highest values across the clusters and bolded values indicate the lowest values across the clusters. Bonferroni post hoc tests were used for pairwise comparison between groups as indicated by superscript numbers, for example where ¹ means significantly different from cluster 1.

Table 2 Demographical information across the clusters

Demographics	Cluster 1		Cluster 2		Cluster 3		P value
	Mean	SD	Mean	SD	Mean	SD	
Age (years)	<u>44</u> ³	13	43	13 ³	36 ^{1,2}	12	7.90 x 10 ⁻²⁴
BMI (kg/m ²)	<u>27.7</u> ^{2,3}	5.3	23.9	3.9 ^{1,3}	25.4 ^{1,2}	4.7	3.04 x 10 ⁻²⁶
W.C. (m)	<u>0.93</u> ^{2,3}	0.14	0.82	0.11 ^{1,3}	0.85 ^{1,2}	0.13	8.48 x 10 ⁻³²
Gender (M/F)	161/165 ³		141/292 ³		266/329 ^{1,2}		3.87 x 10 ⁻⁶
<i>Frequency % (N)</i>							
Germany	19.9 (65)		19.2 (83)		7.9 (47)		1.19 x 10 ⁻⁸
Greece	13.2 (43)		5.5 (24)		20.0 (119)		2.39 x 10 ⁻¹⁰
Ireland	15.6 (51)		20.6 (89)		10.6 (63)		5.34 x 10 ⁻⁵
Netherlands	16.3 (53)		14.1 (61)		14.6 (87)		0.693
Poland	15.0 (49)		17.1 (74)		10.6 (63)		0.008
Spain	8.3 (27)		8.5 (37)		21.3 (127)		1.08 x 10 ⁻¹⁰
United Kingdom	11.7 (38)		15.0 (65)		15.0 (89)		0.325

N, number; W.C., waist circumference. Values are presented as means \pm standard deviations. One-way ANOVA was used to examine the differences across the clusters with exception of gender and country where chi-square was used instead. Underlined values indicate the highest values across the clusters and bolded values indicate the lowest values across the clusters. Bonferroni post hoc tests were used for pairwise comparison between groups as indicated by superscript numbers, for example where ¹ means significantly different from cluster 1.

Table 3 Dietary intakes across the clusters

Nutrient	Cluster 1		Cluster 2		Cluster 3		P value [†]
	Mean	SD	Mean	SD	Mean	SD	
Energy (kJ)	11370	5064	10270	4015	10816	4592	1.00
Total fat (%)	35.9	5.8	36.1	6.3	35.7	5.8	1.00
Saturated fat (%)	14.5	3.2	14.0	3.3	14.0	3.0	1.00
Monounsaturated fat (%)	13.6	3.0	13.7	3.3	13.9	3.1	0.864
Polyunsaturated fat (%)	5.7 ²	1.4	<u>6.1</u> ^{1,3}	1.5	5.6 ²	1.4	0.048
Protein (%)	16.7	3.5	17.1	4.0	17.2	3.5	0.576
Carbohydrate (%)	45.3	7.3	46.1	7.9	46.3	7.4	1.00
Sugars (%)	20.6	6.2	22.1	6.3	20.6	5.3	1.00
Alcohol (%)	<u>4.2</u> ^{2,3}	4.5	2.9 ¹	3.5	3.1 ¹	3.4	1.85 x 10 ⁻³
Salt (g)*	8	4	7	3	7	4	1.00
Fibre (g)*	30 ²	14	<u>32</u> ^{1,3}	15	28 ²	15	9.60 x 10 ⁻⁶
Vitamin A (µg)*	1720 ³	1150	<u>1884</u> ³	1048	1500 ^{1,2}	900	1.98 x 10 ⁻⁵
Vitamin D (µg)*	6 ²	9	<u>8</u> ^{1,3}	18	5 ²	5	3.33 x 10 ⁻⁷
Vitamin E (mg)*	15 ²	13	<u>20</u> ^{1,3}	35	15 ²	21	1.20 x 10 ⁻²
Carotene (µg)*	6243 ²	10534	<u>7313</u> ^{1,3}	5257	5015 ²	3557	1.02 x 10 ⁻⁶
Retinol (µg)*	1340	10175	665	511	664	576	1.00
Thiamin (mg)*	4	10	5	9	4	6	0.600
Riboflavin(mg)*	4	7	4	8	3	6	1.00
Folate (µg)*	424 ²	200	<u>443</u> ^{1,3}	221	405 ²	211	2.26 x 10 ⁻³

Vitamin B6 (mg)*	4	9	5	11	4	11	0.264
Vitamin B12 (µg)*	19	91	19	71	15	67	0.144
Vitamin C (mg)*	219 ²	230	<u>270</u> ^{1,3}	325	192 ²	195	1.48 x 10 ⁻⁸
Calcium (mg)*	1328	656	1261	549	1289	635	1.00
Iron (mg)*	18	11	17	8	17	8	1.00

*Adjusted for energy (kJ). †General linear models were calculated on log transformed values where necessary and adjusted for multiple comparisons. Values are presented as means ± standard deviations. P values provided by general linear models controlling for age, gender, BMI and country where appropriate. Bonferroni post hoc tests used to examine pairwise comparisons between groups with the exception of vitamin D where LSD post hoc tests were used instead. Differences between clusters are indicated by superscript numbers where ¹ means significantly different from cluster 1.

Table 4 Food group intakes across the clusters

Food group (g)	<u>Cluster 1</u>		<u>Cluster 2</u>		<u>Cluster 3</u>		P value*
	Mean	SD	Mean	SD	Mean	SD	
Rice, pasta and grains	81	62	75	58	90	76	0.608
Savouries	24 ²	26	11 ^{1,3}	13	<u>28</u> ²	33	1.27 x 10 ⁻⁴
White bread/rolls/ scones/croissants	55 ²	92	34 ^{1,3}	73	<u>71</u> ²	120	1.36 x 10 ⁻⁵
Wholemeal and brown bread	102	126	100	122	85	156	1.00
Breakfast cereals and porridge	53	82	76	109	48	66	1.00
Biscuits, cakes and pastries	70	133	68	80	70	88	0.512
Wholemilk	45	175	32	93	36	106	1.00
Low fat and skimmed milks	166	217	164	218	184	225	1.00
	196		163		189		
Other milks, milk based beverages and other beverages		314		221		300	1.00
Creams, ice creams and desserts	12	15	9	21	7	10	0.128
Cheese	36	37	37	37	32	35	0.288
Yoghurts	79 ²	96	<u>91</u> ^{1,3}	107	75 ²	128	0.032
Egg and egg dishes	31	38	31	41	32	40	1.00
Butter, fat spreads and hard cooking fats	13	19	9	12	8	11	1.00
Low fat spreads and oils	10	10	9	9	10	11	1.00
Potatoes	60	67	52	48	51	69	1.00
Chips and processed potatoes	25	27	18	21	25	34	1.00
Vegetables and vegetable dishes	190	145	225	194	146	114	0.480

Fruit juices and smoothies	125	176	126	171	114	158	1.00
Fruit	253 ²	216	<u>355</u> ^{1,3}	306	228 ²	189	1.39 x 10 ⁻⁸
Savoury snacks	9	13	9	13	10	14	1.00
Fish, fish dishes and products	55 ²	40	<u>71</u> ^{1,3}	53	66 ²	57	8.16 x 10 ⁻⁴
Red meat	43	37	30	39	41	36	0.704
Poultry	34	35	30	39	35	33	1.00
Meat products	49	51	34	49	45	52	0.832
Red meat dishes	34	58	30	37	33	34	1.00
Alcoholic beverages	211	257	129	180	156	207	0.064
Sugar syrups, preserves and sweeteners	11	14	9	12	10	15	1.00
Confectionary	29	48	21	24	25	30	1.00
Soups, sauces and condiments	100	73	91	72	94	85	1.00
Low energy beverages	556	530	604	509	434	479	0.128
High energy beverages	34	73	12	33	44	169	0.064
Supplement users (%)	37.2 ²		54.3 ^{1,3}		37.5 ²		9.31 x 10 ⁻⁸

*General linear models were calculated on logged values where necessary and adjusted for multiple comparisons. Values are presented as means \pm standard deviations. P values provided by general linear models controlling for age, gender, BMI and country where appropriate. Frequency of supplement users was assessed using chi-squared analysis. Bonferroni post hoc tests used to examine pairwise comparisons between groups as indicated by superscript numbers where ¹ means significantly different from cluster 1.

Table 5 Range of values across the clusters and cut-offs used for the development of the targeted dietary advice

	Cluster 1	Cluster 2	Cluster 3	Cut-offs	
Total cholesterol (mmol/L)	3.987 - 6.033	3.878 - 5.702	3.472 – 5.028	<i>Desirable</i> < 5	<i>High</i> > 5
Total carotenoids (µM)	0.82 – 1.74	1.437 - 2.863	0.807 – 1.613	<i>Desirable</i> > 1.5	<i>Low</i> < 1.5
Total sat fat (%)	High	Low	Medium	N/A	
Omega-3 index (%)	4.232 - 6.088	5.266- 7.854	4.494 - 6.326	<i>Desirable</i> ≥ 8	<i>Low</i> < 4
Dietary fibre (g)	16.21 - 43.29	17.09 – 47.71	13.62 - 42.92	Males	
				18-50 yrs	<i>Desirable</i> ≥ 38 <i>Low</i> < 38
				> 50 yrs	≥ 30 < 30
				Females	
				18-50 yrs	<i>Desirable</i> ≥ 25 <i>Low</i> < 25
				> 50 yrs	≥ 21 < 21
Salt (g)	3.92 - 11.84	3.94 – 10.00	3.48 – 11.44	18-50 yrs	<i>Desirable</i> ≤ 3.75 <i>High</i> > 3.75
				51-70 yrs	≤ 3.25 > 3.25
				> 70 yrs	< 3 > 3
Folate (µg)	224.42 – 623.24	221.88 – 664.04	193.66 – 615.80	<i>Desirable</i> ≥ 320	<i>Low</i> < 320
Calcium (mg)	671.48 – 1984.10	712.24 – 1809.72	654.41 - 1924.15	Males	
				18-70yrs	<i>Desirable</i> ≥ 800 <i>Low</i> < 800
				>70yrs	≥ 1000 < 1000
				Females	
				18-50 yrs	<i>Desirable</i> ≥ 800 <i>Low</i> < 800
				> 50 yrs	≥1000 < 1000

				Males		
				> 18 yrs	<i>Desirable</i> ≥ 6	<i>Low</i> < 6
Iron (mg)	6.94 – 29.22	8.83 – 25.45	9.15 – 24.75	Females		
				18-50 yrs	<i>Desirable</i> ≥ 8.1	<i>Low</i> < 8.1
				> 50y rs	≥ 5	< 5
				<i>Normal</i>	<i>Overweight</i>	<i>Obese</i>
BMI (kg/m ²)	22.43 – 32.93	20.09 – 27.79	20.66 – 30.04	18.5 – 24.99	≥ 25	≥ 30
				Males	<i>Desirable</i> < 102	<i>High</i> ≥ 102
W.C. (m)	0.79 – 1.07	0.71 – 0.93	0.72 – 0.98	Females	< 88	≥ 88

Table 6 Agreement between the proposed targeted dietary advice and the individualised dietary advice method adopted within the Food4Me study

Agreement between targeted and individualised methods (%)	
Cluster 1	83
Cluster 2	74
Cluster 3	88
Total	82

The agreement between the targeted and individualised method is at the level of the delivery of the same dietary message.

Table 7 Number of messages given as per the targeted dietary advice

Number of messages given	No. of times (%)
2	13 (7)
3	46 (26)
4	50 (28)
5	51 (28)
6	20 (11)

This table shows the number of dietary messages given using the proposed targeted dietary advice method.